



The Child with Recurrent Mycobacterial Disease

Brian Reed¹ · William K. Dolen¹

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Abstract

Purpose of Review Many genetic conditions predispose affected individuals to opportunistic infections. A number of immunodeficiency diseases, including genetic defects termed Mendelian susceptibility to mycobacterial disease (MSMD), permit infection from many different strains of mycobacteria that would otherwise not cause disease. These include tuberculous and nontuberculous mycobacteria, and bacille Calmette-Guérin vaccine (BCG). Patients may present with infections from other organisms that depend on macrophage function for containment. Defects in multiple genes in the IL-12 and NFKB signaling pathways can cause the MSMD phenotype, some of which include *IL12RB1*, *IL12B*, *IKBK*, *ISG15*, *IFNGR1*, *IFNGR2*, *CYBB*, *TYK2*, *IRF8*, and *STAT1*.

Recent Findings Multiple autosomal recessive and dominant, and 2 X-linked recessive gene defects resulting in the MSMD phenotype have been reported, and others await discovery. This review presents the known gene defects and describes clinical findings that result from the mutations.

Summary If MSMD is suspected, a careful clinical history and examination and basic immunodeficiency screening tests will narrow the differential diagnosis. A specific diagnosis requires more sophisticated laboratory investigation. Genetic testing permits a definitive diagnosis, permitting genetic counseling. Mild cases respond well to appropriate antibiotic therapy, whereas severe disease may require hematopoietic stem cell transplantation.

Keywords Mendelian susceptibility to mycobacterial disease · MSMD · IL-12 · IFN γ · Salmonella · Candida · Atypical mycobacteria · Mycobacterium avium complex · Bacille Calmette-Guérin · Tuberculosis

Introduction

The mycobacteria are found in soil and water. In humans, they are commensal organisms, the nontuberculous mycobacteria being epithelial colonists of the gastrointestinal, respiratory, and genitourinary systems, as well as the skin. Limbs of the innate and adaptive immune systems normally keep the organisms in check. Either hyporesponsiveness or hyperresponsiveness to these organisms can result in clinical disease [1]. In some countries, bacille Calmette-Guérin (BCG) vaccine, a live vaccine

developed from *Mycobacterium bovis*, is used to prevent tuberculosis. Individuals with certain primary immunodeficiency diseases (including severe combined immunodeficiency, hyper IgM syndrome, and chronic granulomatous disease) can develop disseminated mycobacterial infection following immunization. An interesting series of inherited or sporadic defects that mostly affect the interferon-gamma (IFN γ) pathway have been termed Mendelian susceptibility to mycobacterial disease (MSMD). Nontuberculous mycobacterial disease can also occur in patients with secondary immunodeficiency, including HIV-1 disease, malignancy, immunosuppressive pharmacotherapy, or cystic fibrosis [2•].

The professional phagocytes (neutrophils, dendritic cells, macrophages, monocytes) of the innate immune system are key players, whose pattern recognition receptors (PRRs) can interact with mycobacterial danger-associated molecular patterns (DAMPs). Mycobacterial DAMPs are found in the cell wall. PRRs important in mycobacterial recognition include Toll-like receptors (TLR-2, TLR-4, TLR-6, TLR-9), NOD2, MINCLE and MARCO, Dectin-1, Dectin-2, Dectin-3, DC-SIGN, Galectin-3, CD36, mannose-binding lectin (MBL),

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✉ William K. Dolen
bdolen@augusta.edu

Brian Reed
breed2@augusta.edu

¹ Department of Pediatrics, Allergy-Immunology and Pediatric Rheumatology Division, Medical College of Georgia at Augusta University, 1120 15th Street, Augusta, GA 30912, USA

and NLRP3 [1]. Although there is some functional redundancy in the PRR system, loss of function defects, or absence of innate immune cells bearing them, can increase susceptibility to mycobacterial infection and, in some cases, also infection with *Salmonella* sp., *Candida* sp., viruses, and other organisms. In the adaptive immune system, cytotoxic (CD8+) and helper (CD4+) T cells provide host defense against mycobacterial infection. T cell deficiency, primary or acquired, permits infection. Some MSMD defects are found in the NF κ B signaling pathway. The interleukin-12 (IL-12) signaling pathway, with production of interferon-gamma (IFN γ), is particularly important for host resistance to mycobacteria, and defects in this complex pathway account for most known causes of MSMD. Among other roles in the innate and adaptive immune system, IFN γ activates macrophages, enhancing their ability to kill intracellular organisms, such as mycobacteria and *Salmonella* sp. Other organisms handled by macrophages include *Klebsiella* sp., *Histoplasma* sp., *Paracoccidioides* sp., *Coccidioides* sp., and *Cryptococcus* sp. [3•].

Infection from weakly virulent *Salmonella* sp. and mycobacteria such as nontuberculous mycobacteria and BCG is uncommon to very rare. Some patients have mild disease, whereas others have life-threatening, difficult-to-treat, and sometimes fatal disease. Patients typically present with mycobacterial infections in childhood [4••]. Some adult patients presenting with the MSMD phenotype have autoantibodies to IFN γ [5•] or GATA2 deficiency. Here, we will concisely review the currently known genetic defects that may result in the MSMD phenotype (including some that are not usually classified as part of the MSMD family), discuss clinical features of the defects, following the classification and order given in Table 1, and comment on patient evaluation (Table 2) and therapy.

TLR-2

An important PRR of the innate immune system, Toll-like receptor 2 (TLR-2) recognizes mycobacterial lipoproteins and is responsible for inducing IL-12 production in macrophages. Mutations can increase susceptibility to tuberculosis [9] and leprosy (*M. leprae*, *M. lepromatosis*).

IL-12/23R β 1

Interleukin 12 (IL-12) is produced by dendritic cells, macrophages, and neutrophils in response to activating signals. It promotes differentiation of naïve T cells into Th1 cells and stimulates T cells and NK cells to produce IFN γ and TNF α . Production of IFN γ enhances macrophage phagocytosis, which is important in the host defense against mycobacteria. The receptor is a heterodimer. The IL-12R β 1 subunit binds

with IL-12R β 2 to form the IL-12 receptor, or with IL-23R to form the IL-23 receptor. A defect in the *IL12RB1* gene (which produces IL-12/23R β 1) causes immunodeficiency 30, the most common known cause of MSMD [3•]. Affected patients are susceptible to infection with bacille Calmette-Guérin (BCG), other atypical mycobacteria, *M. tuberculosis*, and *Salmonella* sp. [10•]. Some patients have presented with associated enteropathy and hypogammaglobulinemia [11]. The IL-12R β 1 variation database lists 220 individuals and 192 unique DNA variants [12]. Inheritance is autosomal recessive.

IL-12/23 p40 Subunit

IL-12 is a heterodimer encoded by 2 separate genes, IL12A (p35 subunit) and IL12B (p40 subunit). The IL-12 p40 subunit also heterodimerizes with the IL-23 p19 subunit (IL23A) to form IL-23. Binding of IL-12 to its receptor activates helper T cells and NK cells. A defect in IL12B (IL-12 p40 subunit), causes immunodeficiency 29, another commonly reported cause of MSMD. Patients typically have BCG disease and/or salmonellosis, but tuberculosis and atypical mycobacterial disease have been noted. At least 49 patients have been reported [13•]. Inheritance is autosomal recessive.

IFN γ R1

The receptor for human interferon-gamma (IFN γ), the type II interferon, is a heterodimer of IFN γ R1 (that binds IFN γ) and IFN γ R2. IFN γ activates macrophages, and plasma IFN γ levels can be elevated in patients with IFN γ R1 or IFN γ R2 deficiency [14]. IFN γ R1 deficiency can present in autosomal recessive (AR) and autosomal dominant (AD) forms. AR deficiency (immunodeficiency 27A) presents with severe, early-onset infections with atypical mycobacteria or BCG. Tuberculosis and salmonellosis have been reported. Patients with the AD deficiency (immunodeficiency 27B) have less severe, later onset mycobacterial infections, but are more prone to *Mycobacterium avium* complex osteomyelitis [15, 16].

IFN γ R2

Defects in IFN γ R2 (immunodeficiency 28), which is involved in signal transduction, also result in severe, early-onset mycobacterial infections. In 1998, Dorman and Holland reported a child with disseminated *M. fortuitum* and *M. avium* complex infection and a mutation in the extracellular domain of IFN γ R2 [17]. Other mutations have been reported [18].

Table 1 Genetic causes of recurrent mycobacterial infections, including Mendelian susceptibility to mycobacterial disease. Note that sporadic mutations can occur

| Disorder | Gene | Inheritance | MIM entry (gene) | MIM entry (disease) | Comments |
|--|----------------|-------------|------------------|---------------------|--|
| IL-12 signaling pathway (type II interferon) | | | | | |
| TLR-2 deficiency | <i>TLR2</i> | AD | *603028 | None | TB, leprosy |
| IL-12/23R β 1 deficiency (IMD30) | <i>IL12RB1</i> | AR | *601604 | #614891 | Most common. Elevated IgE; enteropathy, hypogammaglobulinemia; atypical mycobacteria, BCG, <i>Candida</i> , <i>Salmonella</i> , TB |
| IL-12/23 p40 subunit deficiency (IMD29) | <i>IL12B</i> | AR | *161561 | #614890 | BCG, <i>Candida</i> , <i>Salmonella</i> ; TB and atypical mycobacteria also reported |
| IFN γ R1 deficiency AR (IMD27A) | <i>IFNGR1</i> | AR | *107470 | #209950 | Severe, early-onset BCG, atypical mycobacterial infections, TB; <i>Salmonella</i> ; <i>Listeria</i> ; some viruses; elevated plasma IFN γ |
| IFN γ R1 deficiency AD (IMD27B) | <i>IFNGR1</i> | AD | *107470 | #615978 | Less severe mycobacterial, <i>Salmonella</i> disease; <i>M. avium</i> complex osteomyelitis |
| IFN γ R2 deficiency (IMD28) | <i>IFNGR2</i> | AR | *147569 | #614889 | Severe, early-onset BCG, mycobacterial, <i>Salmonella</i> disease; some viruses; elevated plasma IFN γ |
| STAT1 deficiency AD (IMD31A) | <i>STAT1</i> | AD | *600555 | #614892 | Mild disease from BCG, <i>M. avium</i> complex, and TB |
| STAT1 deficiency AR (IMD31B) | <i>STAT1</i> | AR | *600555 | #613796 | Severe BCG, mycobacterial and viral disease |
| JAK1 deficiency | <i>JAK1</i> | AR | *147795 | None | Atypical mycobacterial disease |
| ISG15 deficiency (IMD38) | <i>ISG15</i> | AR | *147571 | #616126 | Severe BCG, mycobacterial disease; intracranial calcification |
| RORC deficiency (IMD42) | <i>RORC</i> | AR | *602943 | #616622 | BCG, mycobacterial, and candida disease |
| GATA2 deficiency/MonoMAC (IMD21) | <i>GATA2</i> | AD | *137295 | #614172 | Mostly in adults |
| SLC11A1/NRAMP deficiency | <i>SLC11A1</i> | (Autosomal) | *600266 | None | TB, leprosy, rheumatoid arthritis |
| Type I interferon | | | | | |
| IRF8 deficiency AD (IMD32A) | <i>IRF8</i> | AD | *601565 | #614893 | Disseminated BCG |
| IRF8 deficiency AR (IMD32B) | <i>IRF8</i> | AR | *601565 | #614894 | Disseminated BCG, candidiasis |
| TYK2 deficiency (IMD35) | <i>TYK2</i> | AR | *176941 | #611521 | Hyper IgE (not always); BCG, mycobacteria, <i>Salmonella</i> , viruses |
| NFKB signaling pathway | | | | | |
| NFKBIA deficiency | <i>NFKBIA</i> | AD | *164008 | #612132 | Anhidrotic ectodermal dysplasia; BCG, mycobacterial disease |
| IKBKB/IKK β deficiency (IMD15) | <i>IKBKB</i> | AR | *603258 | #615592 | Early-onset severe viral, fungal, bacterial infections, mycobacterial disease |
| IKBKG/IKK γ /NEMO | <i>IKBKG</i> | XLR | *300248 | #300636 | <i>M. avium</i> complex; conical teeth reported |
| CARMIL2/RLTPR | <i>CARMIL2</i> | AR | *610859 | None | TB; candidiasis |
| Other | | | | | |
| CYBB deficiency (IMD34) | <i>CYBB</i> | XLR | *300481 | #300645 | BCG disease, TB; the DHR respiratory burst assay may be normal in these patients [6••] |

MIM numbers are from Online Mendelian Inheritance in Man (OMIM) [7]. Adapted from Picard et al. [8•]

AD autosomal dominant, AR autosomal recessive, BCG bacille Calmette-Guérin, DHR dihydrorhodamine, IMD immunodeficiency, TB tuberculosis, XLR x-linked recessive

STAT1

Signal from the IFN γ R, as well as the type I interferon receptor, is transduced via a JAK/STAT pathway that uses JAK1

(Janus kinase 1) and JAK2, and STAT1 (signal transducer and activator of transcription 1). JAK phosphorylation of STAT1 produces a homodimer that binds to DNA sequences in the nuclear gamma interferon-activated site. STAT1 deficiency

Table 2 Diagnostic considerations of the child with recurrent mycobacterial infections

| Investigation | Comments |
|---|--|
| Detailed medical history | Immunosuppressive pharmacotherapy? Consanguinity? Consider malignancy, cystic fibrosis |
| Detailed physical examination | |
| CBC with differential | Absolute lymphocyte count (SCID) Absolute monocyte count (GATA2 deficiency) |
| IgG, A, M, E levels | |
| Basic lymphocyte cytofluorometry (T, B, NK) | Evaluate for SCID |
| HIV-1 testing | |
| Lymphocyte functional testing | |
| Plasma IFN γ level | Elevated in patients with IFN γ R deficiency |
| QuantiFERON-TB Gold (Qiagen, Hilden, Germany) | Undetectable or low IFN γ production suggests anti-IFN γ antibodies [6••] |
| Genetic testing | Commercial testing panels available; may require DNA sequencing |
| Other laboratory testing | Detailed in a recent review by Esteve-Sole et al. [6••] |

can present in autosomal dominant (immunodeficiency 31A) and autosomal recessive (immunodeficiency 31B) inheritance patterns. The AD form affects IFN γ R signal transduction, producing relatively mild disease from BCG, *M. avium* complex, and TB. The AR form also affects signal transduction from the type I interferon receptor, producing severe mycobacterial and viral disease.

JAK1 (OMIM 147795)

Janus kinase 1 (JAK1) is a tyrosine kinase that adds a phosphate group to a tyrosine amino acid on substrate proteins. JAK1 is required for interferon type I (IFN α , IFN β), and type II (IFN γ) signal transduction. A JAK1 deficiency resulted in atypical mycobacterial disease in a 22-year-old Pakistani male from consanguineous parents [19]. Inheritance is autosomal recessive.

ISG15

ISG15 is a ubiquitin-like modifier induced by interferon α and interferon β , resulting in conjugation to many cellular proteins, and release from monocytes to cause NK cell activation and cytotoxicity [20]. In the absence of ISG15, the amount of IFN γ production is reduced, which may account for the increased susceptibility to mycobacterial disease, as found in immunodeficiency 38. Bogunovic et al. reported 3 patients who developed severe BCG and mycobacterial disease [21]. Intracranial calcification is also reported [22]. Inheritance is autosomal recessive.

RORC

RAR-related orphan receptor C (RORC) encodes for the transcription factor ROR γ . The ROR γ t isoform promotes differentiation of thymocytes into Th17 cells. Homozygous mutations of RORC were discovered in 3 patients from different consanguineous families. RORC deficiency causes immunodeficiency 42, which presents in infancy with candida and mycobacterial infections. Patients have impaired production of IL-17A, IL-17F, and IL-22, and an apparent defect in IFN γ production in response to BCG and IL-12 [23]. Inheritance is autosomal recessive.

GATA2 (“MonoMAC Syndrome”)

GATA-binding protein 2 (GATA2) gene mutations produce immunodeficiency 21, which is associated with a marked decrease (or absence) of monocytes, B cells, NK cells, and dendritic cells. Patients are usually adults, susceptible to various viral, fungal, and mycobacterial infections [24–27].

SLC11A1/NRAMP1

Solute carrier family 11, member 1 (SLC11A1), also known as natural resistance-associated macrophage protein 1 (NRAMP1), is a macrophage membrane protein that also appears to be involved in neutrophil function. Deficiency has been reported to confer susceptibility to tuberculosis and leprosy, as well as rheumatoid arthritis [28].

IRF8

Interferon regulatory factor 8 (IRF8) is a member of a family of transcription factors that regulate type I interferon gene expression [29]. It is only present in immune cells [30]. Deficiency can present as autosomal dominant immunodeficiency 32A, as found in two individuals of Italian descent with disseminated BCG disease and a loss of a dendritic cell population. An autosomal recessive form, termed immunodeficiency 32B, causes monocyte and dendritic cell deficiency with resulting severe infections, including disseminated BCG and candidiasis [31].

TYK2

Tyrosine kinase 2 (TYK2) stabilizes cell surface expression of the interferon α and β receptor subunit 1 (IFNAR1) [32]. TYK2 deficiency produces immunodeficiency 35, reported in a 22-year-old Japanese male with hyper IgE syndrome diagnosed at age 22 months with a BCG infection [33]. He had a history of infections with viruses, fungi, and mycobacteria, and *Salmonella sp.* gastroenteritis at age 15. His healthy, consanguineous parents were heterozygous for the mutation. A few other patients have been reported, and elevated IgE is not a consistent finding. Inheritance is autosomal recessive.

NFKBIA/I κ B α

Nuclear factor of kappa light polypeptide gene enhancer in B cells inhibitor, alpha (NFKBIA, or I κ B α), inhibits the NF κ B complex (along with NFKBIB), inactivating the transcription factor NF- κ B (NFKB) by preventing nuclear translocation [34]. NFKBIA gain of function mutations decreases the production of cytokines and interferons, producing anhidrotic ectodermal dysplasia and susceptibility to various kinds of infections, including a few recently reported patients with BCG and mycobacterial disease [35–37]. Inheritance is autosomal dominant (chromosome 14).

IKBKB/IKK β

Inhibitor of kappa light chain gene enhancer in B cells, kinase of, beta (IKBKB/IKK β /IKK2) and IKBKA/IKK α /IKK1, is a subunit of I κ B kinase (IKK), which phosphorylates I κ B proteins and permits activation of NF- κ B (NFKB). IKBKB and IKBKA associate with IKBKG (NEMO) to form IKK. Mutations in IKBKB cause immunodeficiency 15, which presents in early life with severe

viral, fungal, and bacterial infections, as well as MSMD. Immunoglobulin levels are low, although T and B cells are present. Burns et al. presented a female, born to consanguineous parents, diagnosed with a generalized rash with disseminated BCG at age 3 months. PCR was positive for *M. tuberculosis* complex. Exome sequencing revealed an IKBKB gene mutation [38]. Other mutations have been reported [39].

IKBKG/IKK γ /NEMO

Inhibitor of kappa light chain gene enhancer in B cells, kinase of, gamma (IKBKG, IKK γ), also termed NF- κ B essential modulator (NEMO), is a subunit of I κ B kinase (IKK). Mutations in NEMO can present with varied features, including immunodeficiency 33, X-linked recessive susceptibility to mycobacterial disease. This is probably due to defective IL-12 production, causing low levels of IFN γ . Patients typically present with *M. avium* complex disease, and some have dental abnormalities, including conical teeth. [40, 41].

CARMIL2/RLTPR

RGD-, leucine-rich repeat-, tropomodulin domain-, and proline-rich domain-containing protein (RLTPR), also known as capping protein regulator and myosin 1 linker 2 (CARMIL2), is vital for T cell CD28 co-stimulation in T cells, acting in a scaffolding role to bridge CD28 to CARD11 and the NF κ B pathway. Six patients with CARMIL2 gene mutations were reported by Wang et al.; two of which developed mycobacterial infections. At age 8, 1 patient was diagnosed with multifocal tuberculosis within the cervical lymph nodes, respiratory tract, and digestive tract. The other patient developed miliary tuberculosis at 9 years old. Mucocutaneous candidiasis was also reported [42].

CYBB

Neutrophils kill ingested organisms by using NADPH-oxidase, which assembled from several subunits. One subunit, cytochrome b(-245) or p91-phox, is a heterodimer of a beta subunit (CYBB) and an alpha subunit (CYBA). Deficiency of either subunit can cause chronic granulomatous disease (CGD), but CYBB deficiency is not uniformly associated with a defective neutrophil oxidative burst assay. A CYBB gene defect causes immunodeficiency 34 without CGD features as described by Bustamante et al., who presented a group of 7 male patients with CYBB missense mutations leading to

recurrent BCG disease or recurrent tuberculosis [43]. Inheritance is x-linked recessive.

Conclusions

Infection with BCG or nontuberculous mycobacteria in a child is a rare event that should prompt a diligent search for an underlying immunodeficiency, particularly if there have also been infections with *Salmonella* sp. or *Candida* sp. The list of known causes of MSMD (Table 1) is daunting, and other gene defects await discovery, so evaluating a patient with recurrent mycobacterial infections is a challenging task [6••]. The differential diagnosis can be narrowed somewhat by a detailed history and examination, as well as some basic laboratory tests (Table 2). Because many defects are inherited, a detailed family history (including careful questioning to determine consanguinity) is warranted. Routine laboratory screening tests, such as a CBC with differential, immunoglobulin levels (G, A, M, E), lymphocyte cytofluorometry (T, B, NK), and HIV1 testing, warrant consideration. Lymphocyte functional testing may be indicated.

An absence or marked reduction in monocytes numbers on routine CBC with differential suggests GATA2 deficiency (MonoMAC). Elevation of plasma IFN γ suggests a deficiency in a component of the IFN γ receptor heterodimer [14]. Some forms of specialized testing, including genetic testing panels, are available from commercial reference laboratories, but detailed testing is most effectively coordinated by a primary immunodeficiency center with expert immunology and genetics laboratories [6••]. In some cases, some form of DNA sequencing (such as whole genome sequencing or whole exome sequencing) will be indicated, since new gene defects await discovery. Genetic confirmation of the diagnosis is essential, not only for accurate diagnosis but also for effective genetic counseling. In countries that offer BCG immunization, genetic testing of family members of MSMD patients would be warranted prior to immunization.

Some of the milder forms of MSMD require no treatment other than appropriate antibiotics. Recombinant interferon-gamma injections have also been used in patients with defects that affect IFN γ production, although the product does not have FDA approval for this indication. The more severe forms have been treated with hematopoietic stem cell transplantation. Gene therapy and other approaches are on the horizon [44•].

Compliance with Ethical Standards

Conflict of Interest Drs. Reed and Dolen declare no conflict of interest.

Human and Animal Rights and Informed Consent This article does not contain any studies with human or animal subjects performed by any of the authors.

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